

SEQUENCE LISTING

<110> Cottingham, Ian R.
McCreath, Graham E.

<120> Fusion Proteins Incorporating Lysozyme

<130> 0623.0730002/EKS/BJD

<140> US (to be assigned)

<141> 2001-12-21

<150> US (to be assigned)

<151> 2001-12-21

<150> PCT/GB00/02459

<151> 2000-06-23

<150> GB 9914733.2

<151> 1999-06-23

<150> US 60/147,819

<151> 1999-08-10

<160> 11

<170> PatentIn Ver. 2.1

<210> 1

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<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Linker

<400> 1

Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
1 5 10 15

<210> 2

<211> 4

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Recognition
site for enzymatic cleavage

<400> 2

Ile Glu Gly Arg
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<210> 3

<211> 4

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Recognition

site for enzymatic cleavage

<400> 3

Asp Asp Asp Lys

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<210> 4

<211> 12061

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: DNA sequence of pCLYSM, excluding the bacterial plasmid

<400> 4

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<210> 5

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cleavage site recognised by enterokinase

<400> 5

Phe Pro Thr Asp Asp Asp Lys

1

5

<210> 6

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Linker arm

<400> 6

Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser Ala

1

5

10

15

Ser

<210> 7

<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Enterokinase
cleavage site

<400> 7
Asp Asp Asp Asp Lys
1 5

<210> 8
<211> 15
<212> DNA
<213> Artificial Sequence

<220>
<221> CDS
<222> (1)..(15)

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<223> Description of Artificial Sequence: Normal
lysozyme C-terminal

<400> 8
ggt tgt gga gtg taa
Gly Cys Gly Val
1 5

15

<210> 9
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Normal
lysozyme C-terminal

<400> 9
Gly Cys Gly Val
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<210> 10
<211> 167
<212> DNA
<213> Artificial Sequence

<220>
<221> CDS
<222> (1)..(162)

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<223> Description of Artificial Sequence: C terminal
extension

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 Leu Glu Gly Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Gly Gly
 1 5 10 15
 agc gct agc atg tgc tcc aac ctg tcc acc tgc gtg ctg ggc aag ctg 96
 Ser Ala Ser Met Cys Ser Asn Leu Ser Thr Cys Val Leu Gly Lys Leu
 20 25 30
 agc cag gag ctg cac aag ctg cag acc tac cct agg acc aac acc ggc 144
 Ser Gln Glu Leu His Lys Leu Gln Thr Tyr Pro Arg Thr Asn Thr Gly
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 agc ggc acc cct gga taa tcgat 167
 Ser Gly Thr Pro Gly
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<210> 11
 <211> 53
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: C terminal
 extension

<400> 11
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 1 5 10 15
 Ser Ala Ser Met Cys Ser Asn Leu Ser Thr Cys Val Leu Gly Lys Leu
 20 25 30
 Ser Gln Glu Leu His Lys Leu Gln Thr Tyr Pro Arg Thr Asn Thr Gly
 35 40 45
 Ser Gly Thr Pro Gly
 50